

BIO392-cnv-freq-Exercise

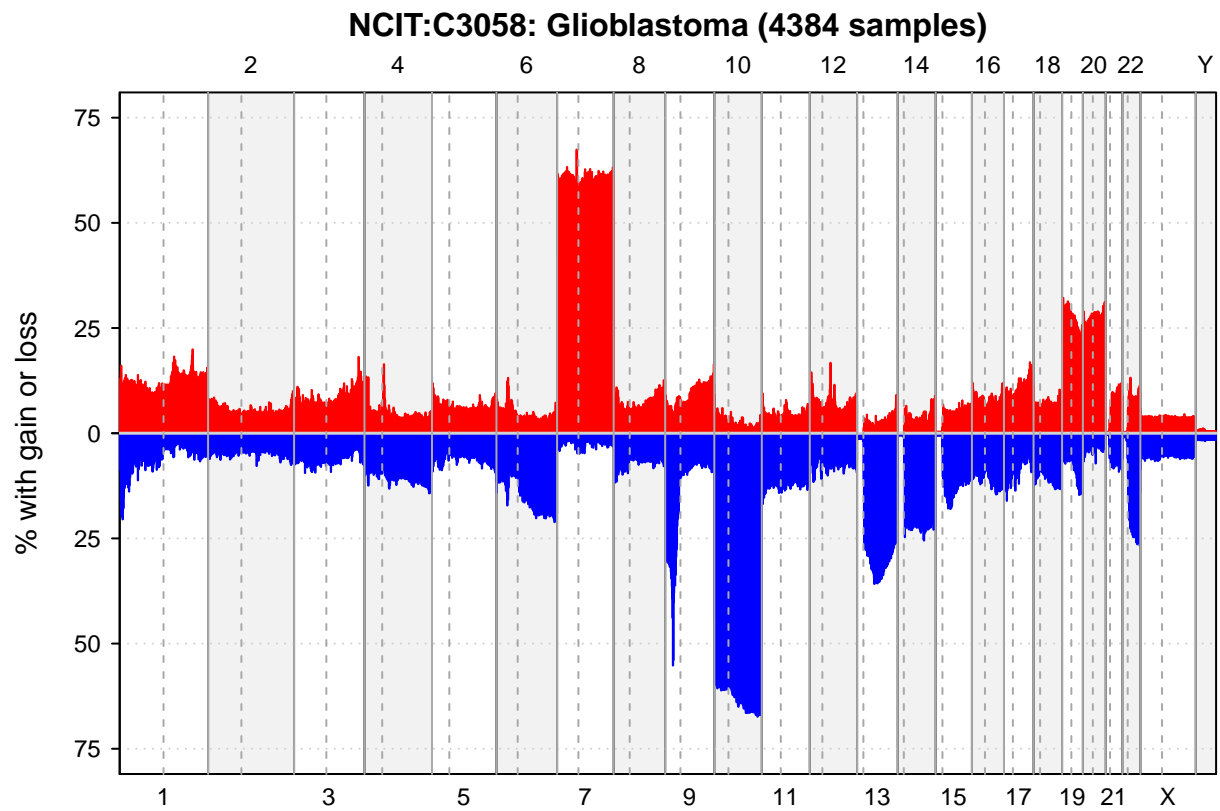
Rebekka Fülleman

2022-09-28

Visualize data

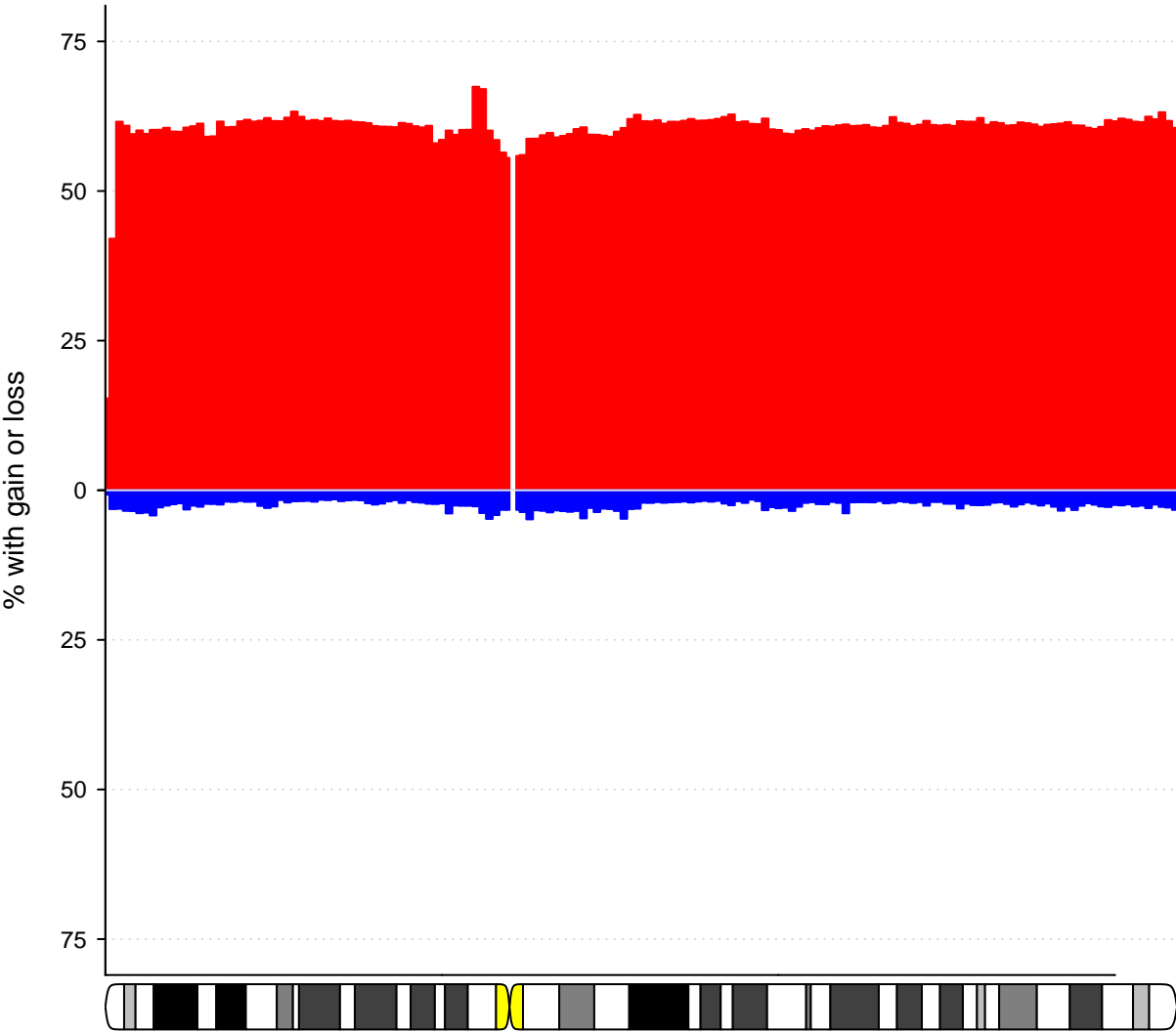
By genome

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pgxFreqplot(freq)
```

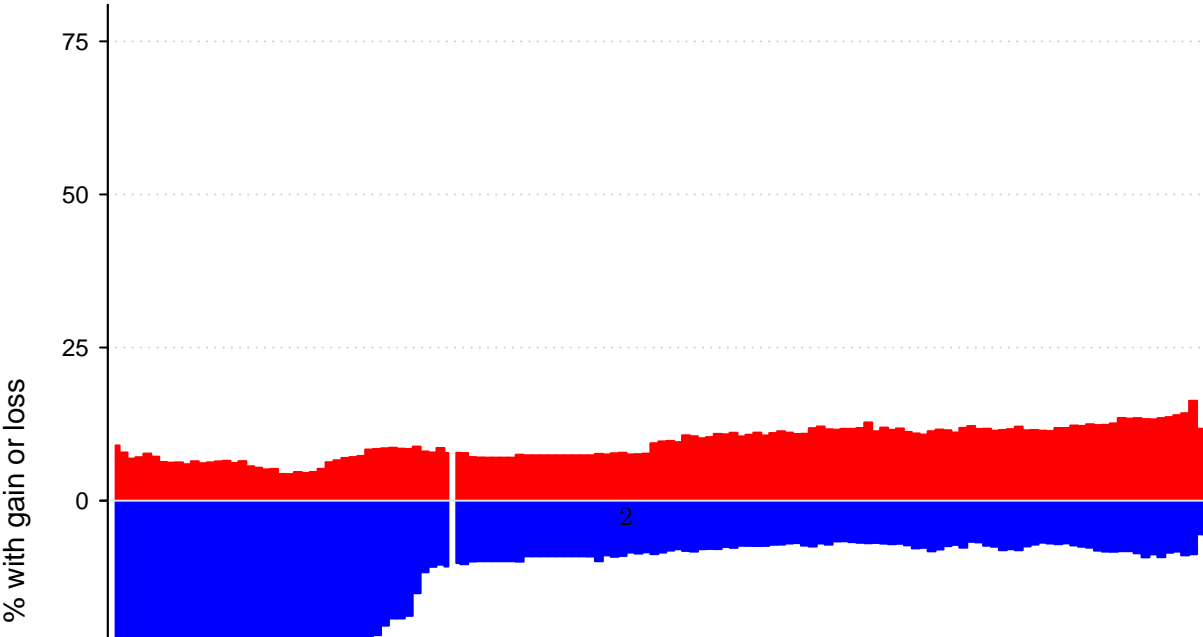


By chromosome

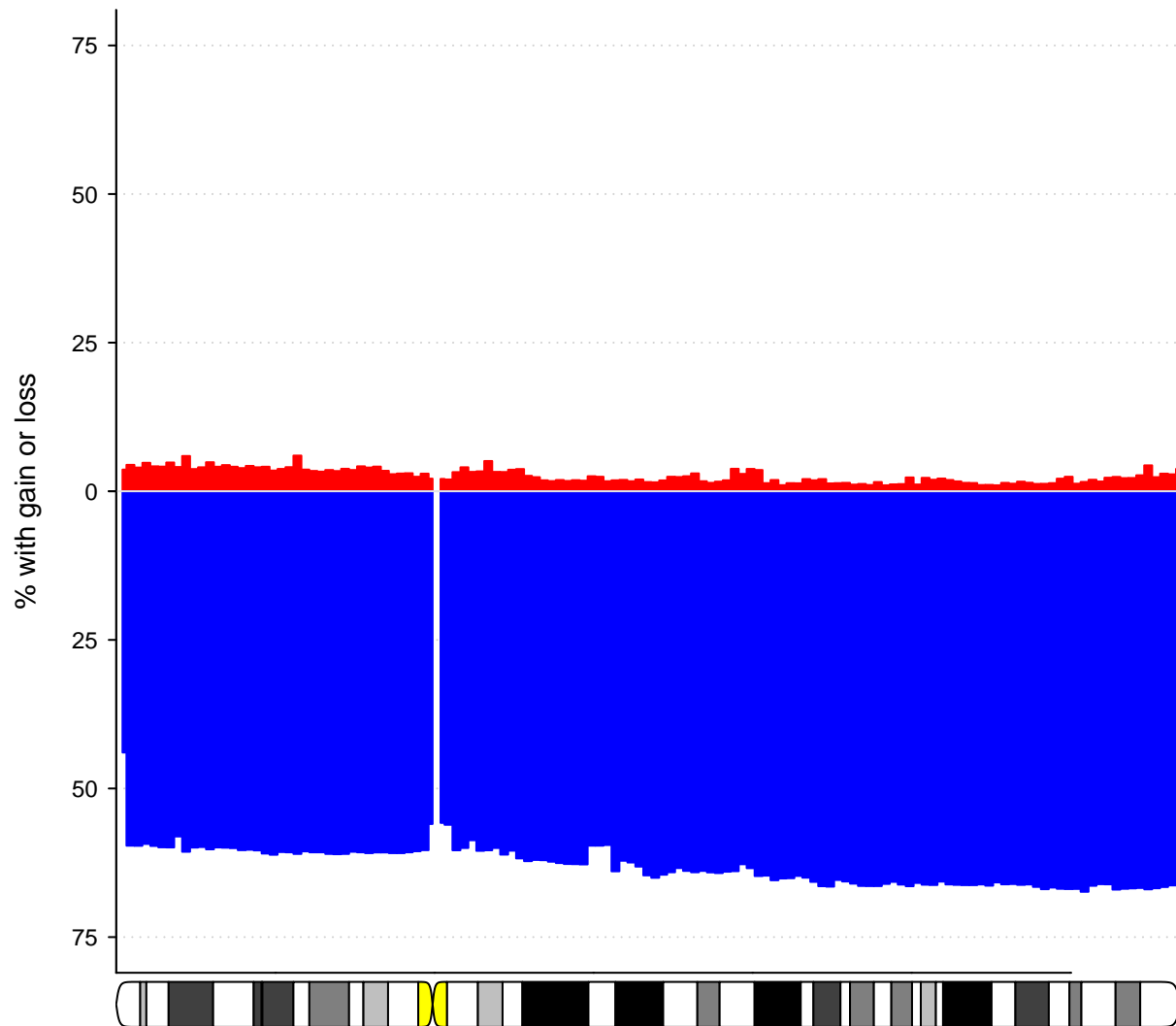
NCIT:C3058: Glioblastoma (4384 samples)
Chromosome 7



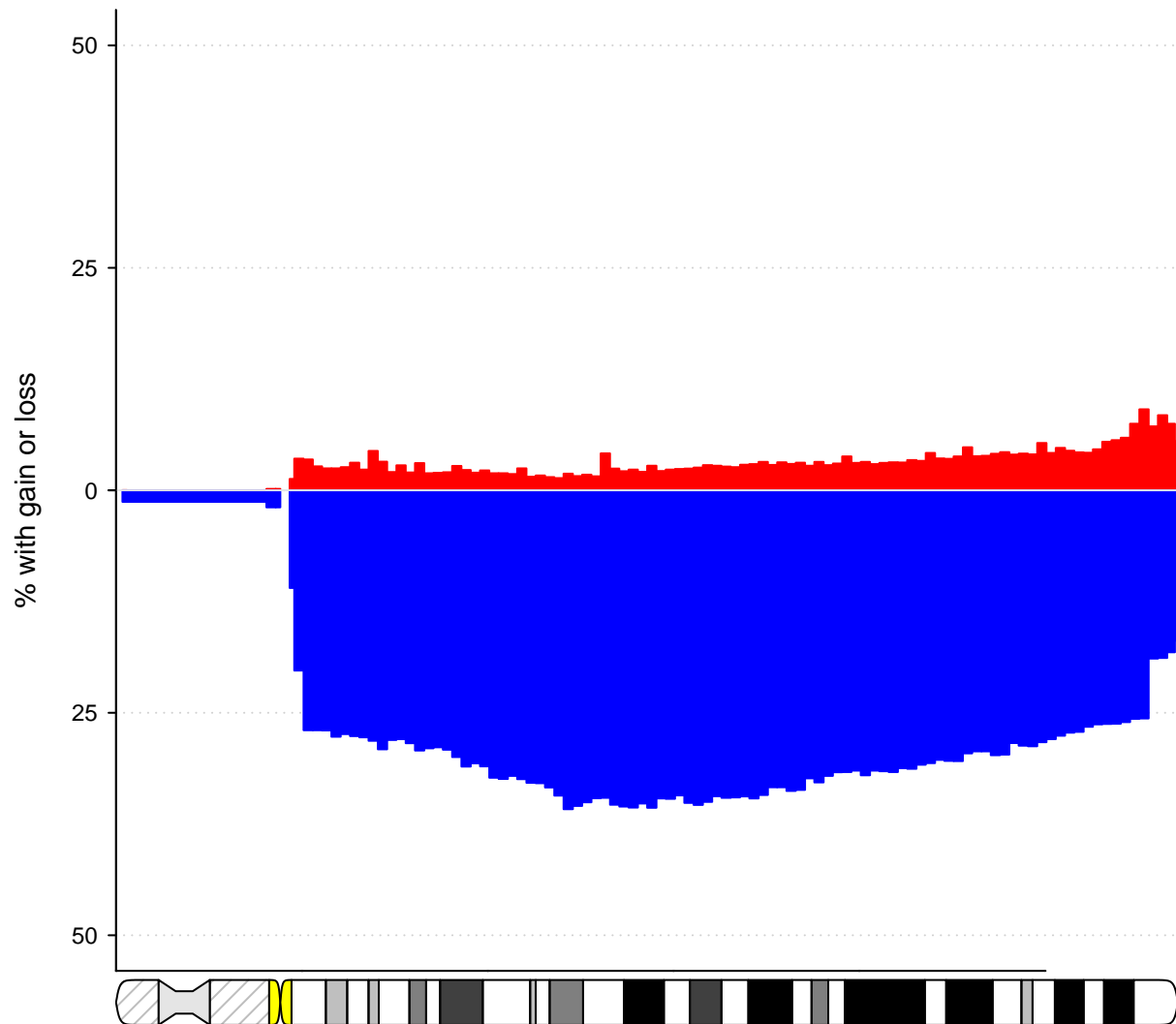
NCIT:C3058: Glioblastoma (4384 samples)
Chromosome 9



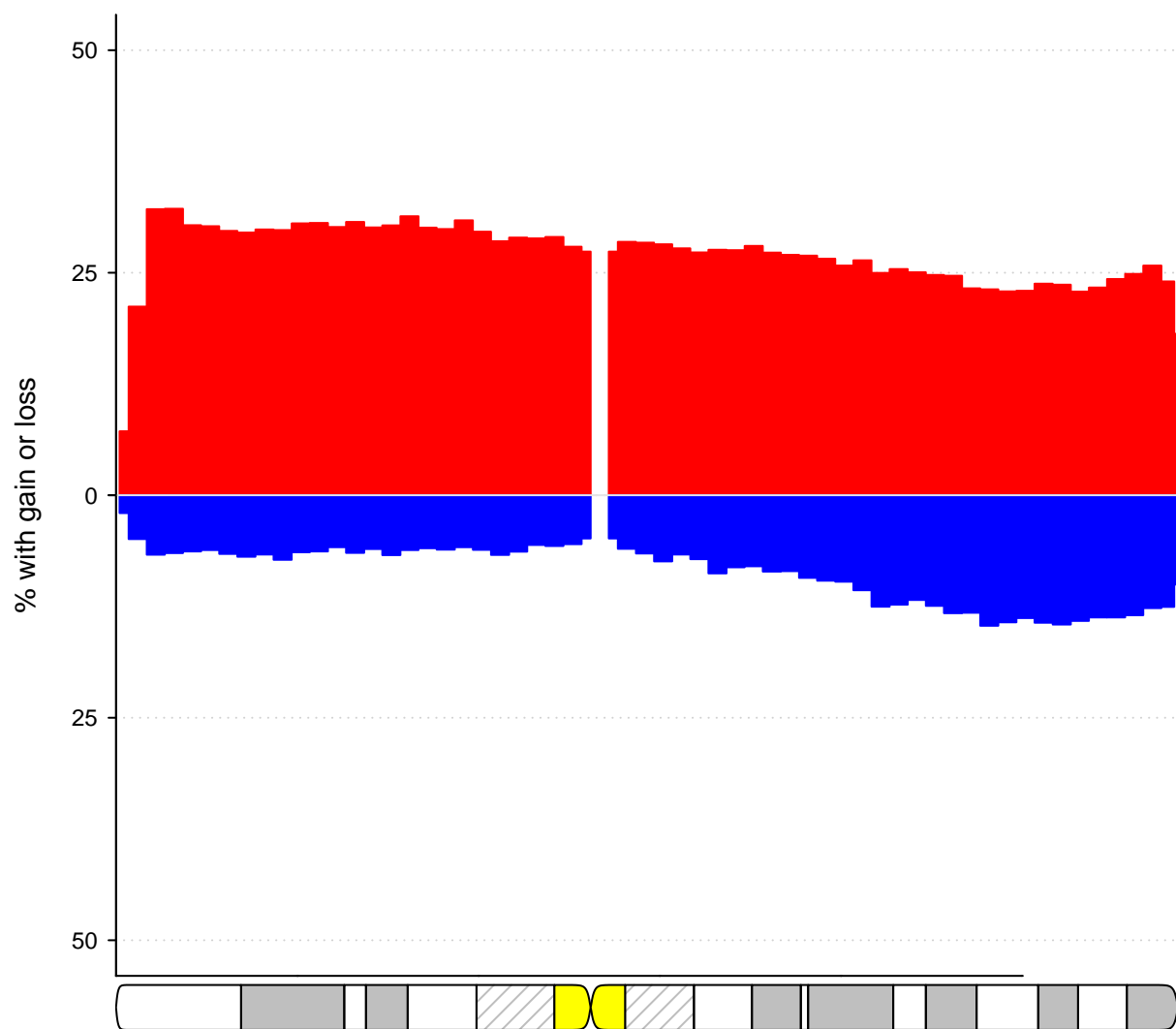
Chromosome 10



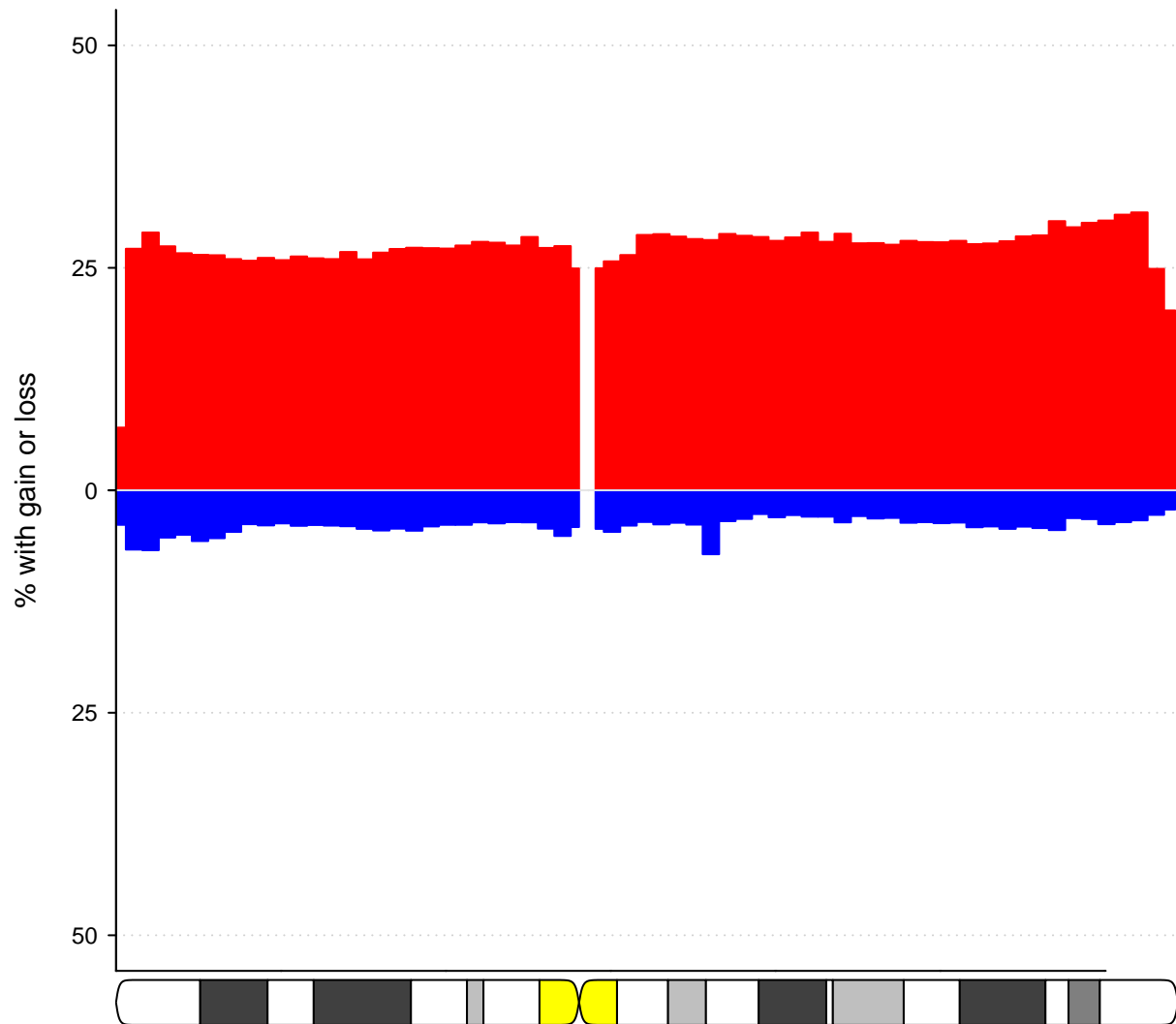
NCIT:C3058: Glioblastoma (4384 samples)
Chromosome 13



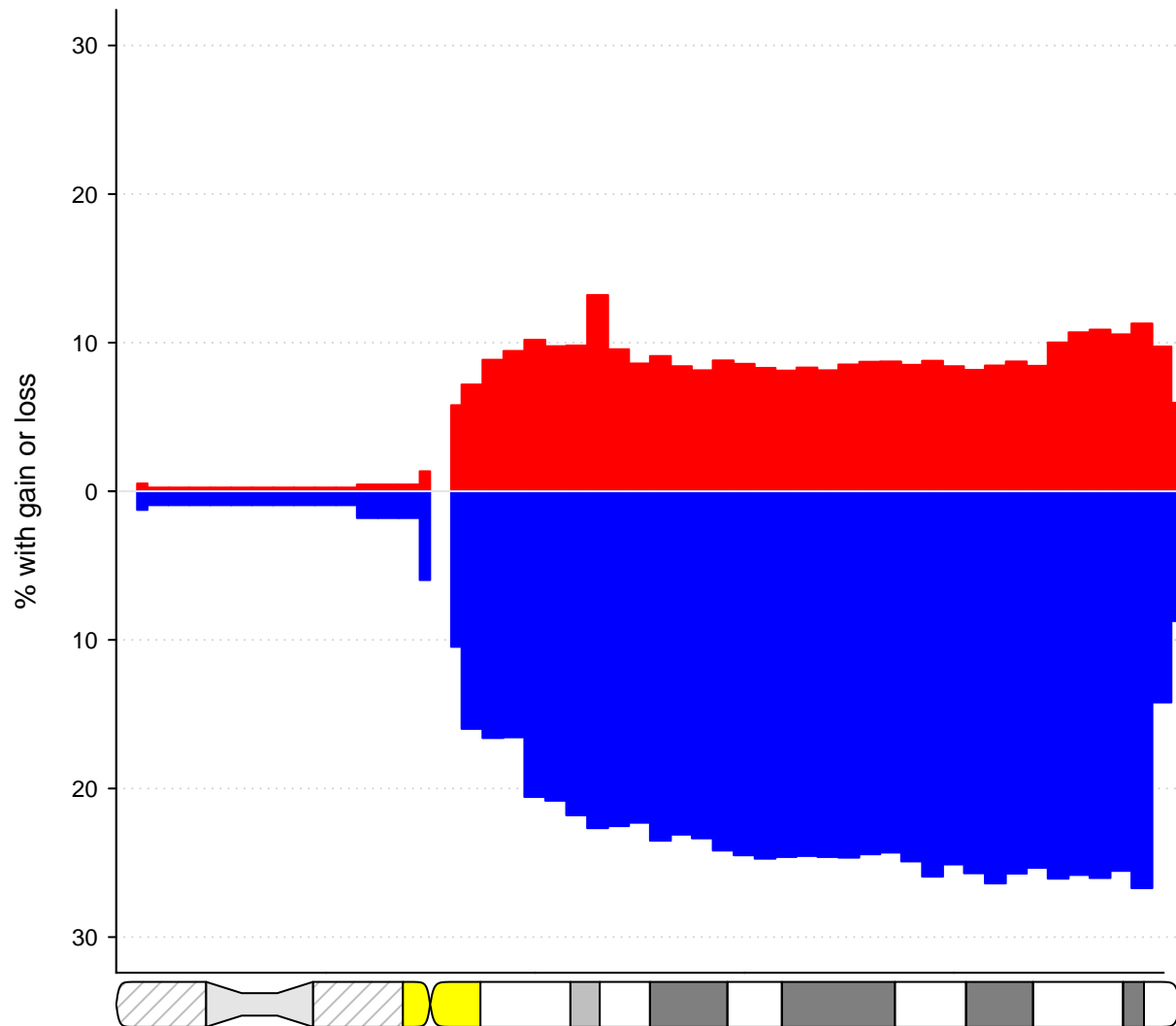
NCIT:C3058: Glioblastoma (4384 samples)
Chromosome 19



NCIT:C3058: Glioblastoma (4384 samples)
Chromosome 20



NCIT:C3058: Glioblastoma (4384 samples)
Chromosome 22



Step5: Analyse the data

According the plot, we can see frequent gains on chromosome 7p, 7q, 19p, (19q,) 20p, 20q and frequent losses on chromosome 9p,10p, 10q, 13q, 22q.